



HA

1

SEQUENCE LISTING

<110> Feussner, Ivo
Hornung, Ellen
Fritsche, Kathrin
Peitzsch, Nicola

<120> Fatty acid desaturase gene from plants

<130> 0050/50669

<140> US 09/721,756

<150> PCT/EP/00/08222

<151> 2000-08-23

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gtc gat cca ccg ttc acg tta agc gat ctg aag aaa gcg att cct acc 152
Val Asp Pro Pro Phe Thr Leu Ser Asp Leu Lys Lys Ala Ile Pro Thr
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His Cys Phe Glu Arg Ser Val Ile Arg Ser Ser Tyr Tyr Val Val His
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Asp Leu Ile Val Ala Tyr Val Phe Tyr Tyr Leu Ala Asn Thr Tyr Ile
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cct ctt att cct aca cct ctg gct tac cta gca tgg ccc gtt tac tgg 296
Pro Leu Ile Pro Thr Pro Leu Ala Tyr Leu Ala Trp Pro Val Tyr Trp
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Tyr	Ser	His	Arg	Asn	His	His	Ala	Asn	Thr	Asn	Ser	Leu	Asp	Asn	Asp		
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Glu	Val	Tyr	Ile	Pro	Lys	Arg	Lys	Ser	Lys	Val	Lys	Ile	Tyr	Ser	Lys		
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Thr	Leu	Gly	Phe	Pro	Leu	Tyr	Leu	Leu	Thr	Asn	Ile	Ser	Gly	Lys	Lys		
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gac agg act cca att ttc aaa gca atg tat aga gag gct aag gaa tgc 1112
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atc tac atc gag ccc gat gag gat agc gag cac aaa ggt gtg ttc tgg 1160
 Ile Tyr Ile Glu Pro Asp Glu Asp Ser Glu His Lys Gly Val Phe Trp
 360 365 370

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 35 40 45

Tyr Tyr Val Val His Asp Leu Ile Val Ala Tyr Val Phe Tyr Tyr Leu
 50 55 60

Ala Asn Thr Tyr Ile Pro Leu Ile Pro Thr Pro Leu Ala Tyr Leu Ala
 65 70 75 80

Trp Pro Val Tyr Trp Phe Cys Gln Ala Ser Ile Leu Thr Gly Leu Trp
 85 90 95

Val Ile Gly His Glu Cys Gly His His Ala Phe Ser Asp Tyr Gln Leu
 100 105 110

Ile Asp Asp Ile Val Gly Phe Val Leu His Ser Ala Leu Leu Thr Pro
 115 120 125

Tyr Phe Ser Trp Lys Tyr Ser His Arg Asn His His Ala Asn Thr Asn
 130 135 140

Ser Leu Asp Asn Asp Glu Val Tyr Ile Pro Lys Arg Lys Ser Lys Val
 145 150 155 160

Lys Ile Tyr Ser Lys Leu Leu Asn Asn Pro Pro Gly Arg Val Phe Thr
 165 170 175
 Leu Val Phe Arg Leu Thr Leu Gly Phe Pro Leu Tyr Leu Leu Thr Asn
 180 185 190
 Ile Ser Gly Lys Lys Tyr Gly Arg Phe Ala Asn His Phe Asp Pro Met
 195 200 205
 Ser Pro Ile Phe Asn Asp Arg Glu Arg Val Gln Val Leu Leu Ser Asp
 210 215 220
 Phe Gly Leu Leu Ala Val Phe Tyr Ala Ile Lys Leu Leu Val Ala Ala
 225 230 235 240
 Lys Gly Ala Ala Trp Val Ile Asn Met Tyr Ala Ile Pro Val Leu Gly
 245 250 255
 Val Ser Val Phe Phe Val Leu Ile Thr Tyr Leu His His Thr His Leu
 260 265 270
 Ser Leu Pro His Tyr Asp Ser Thr Glu Trp Asn Trp Ile Lys Gly Ala
 275 280 285
 Leu Ser Thr Ile Asp Arg Asp Phe Gly Phe Leu Asn Arg Val Phe His
 290 295 300
 Asp Val Thr His Thr His Val Leu His His Leu Ile Ser Tyr Ile Pro
 305 310 315 320
 His Tyr His Ala Lys Glu Ala Arg Asp Ala Ile Lys Pro Val Leu Gly
 325 330 335
 Glu Tyr Tyr Lys Ile Asp Arg Thr Pro Ile Phe Lys Ala Met Tyr Arg
 340 345 350
 Glu Ala Lys Glu Cys Ile Tyr Ile Glu Pro Asp Glu Asp Ser Glu His
 355 360 365
 Lys Gly Val Phe Trp Tyr His Lys Met
 370 375